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Listing of Claims.

Please amend the claims as shown below. This listing of claims will replace all prior versions and listings of the claims in this application.

15. (Previously presented) A method for determining the presence of EBV-positive nasopharyngeal carcinoma or gastric carcinoma cells in a sample of an individual suspected of or at risk for carrying an EBV associated disease, comprising:

(a) amplifying one or more targets from the BKRF1 reading frame spanning nucleotides 107950 – 109872 of EBNA-1,

(b) amplifying one or more target sequence(s) selected from the BARF1 reading frame spanning nucleotides 165504 – 166166,

(c) detecting the presence or absence of the amplified target sequences of steps (a) and (b), and

(d) determining the presence of EBV-positive nasopharyngeal carcinoma or gastric carcinoma cells from the presence of the amplified target sequences of steps (a) and (b).

16. (Previously presented) The method according to claim 15 wherein

-the step of amplifying the BKRF1 reading frame in step (a) is performed using a pair of oligonucleotides, each oligonucleotide comprising a sequence specific for **EBNA-1**, the EBNA-1 specific sequences consisting of

5'-CTCCCTTTACAACCTAAGGC-3' [SEQ.ID.NO.: 2], and

5'-AGAGACAAGGTCCTTAATCGCATCC-3' [SEQ.ID.NO.: 3],

wherein the latter oligonucleotide further comprises a T7

polymerase promoter sequence 5'-aattctaatacgactcactataggg-3' (SEQ ID NO:37);

and

and a pair of oligonucleotides specific for **EBER-1** consisting of

1.1, 5'-CGGGCGGACCAGCTGTACTTGA-3' [SEQ.ID.NO.: 6] provided with a

T7 polymerase promoter sequence 5'-aattctaatacgactcactataggg-3', and

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- the step of amplifying the BARF1 reading frame in step (b) is performed using a pair of oligonucleotides, each oligonucleotide comprising a sequence specific for **BARF-1**, the BARF-1 specific sequences consisting of 5'-GGCTGTCACCGCTTTCTTGG-3' [SEQ.ID.NO.: 23], and 5'-AGTGTGGCACTTCTGTGG-3' [SEQ.ID.NO.: 24], wherein the latter oligonucleotide further comprises a T7 polymerase promoter sequence 5'-aattctaatacgactcactataggg-3' (SEQ ID NO:37).

17. (Previously presented) The method according to claim 15, wherein the RNA is amplified using a transcription based amplification technique.

18. (Previously presented) The method according to claim 17, wherein said amplification technique is NASBA.

Claims 19-26. (Canceled)

27. (Currently amended) The [A] method according to Claim 15, further comprising detecting the presence or absence of the amplified target sequence of (a).

Claims 28-34. (Canceled)